## FIG. 1A

1  "	51 SP.PTVSCQA NNFSSVPLSL PPSTQRLFLQ NNLIRTLRPG TFGSNLLT AP.MTVSCQA HNFAAIPEGI PEDSERIFLQ NNRITFLQQG HFSPAMVT EPKVTTSCPQ QGLQAVPVGI PAASQRIFLH GNRISHVPAA SFRACRNLTI -PT-SCP PR-FLN-IF	101 LWLFSNNLST IYPGTFRHLQ ALEELDLGDN RHLRSLEPDT FQGLERLQSL LWIYSNNITF IAPNTFEGFV HLEELDLGDN RQLRTLAPET FQGLVKLHAL LWLHSNVLAR IDAAAFTGLA LLEQLDLSDN AQLRSVDPAT FHGLGRLHTL LWSN IFLE-LDL-DNLRP-T F-GL-L-L	151 HLYRCQLSSL PGNIFRGLVS LQYLYLQENS LLHLQDDLFA DLANLSHLFL YLYKCGLSAL PAGIFGGLHS LQYLYLQDNH IEYLQDDIFV DLVNLSHLFL HLDRCGLQEL GPGLFRGLAA LQYLYLQDNA LQALPDDTFR DLGNLTHLFL -LC-LLF-GL LQYLYLQ-NL-DD-F- DL-NL-HLFL	201 HGNRLRLLTE HVFRGLGSLD RLLLHGNRLQ GVHRAAFRGL SRLTILYLFN HGNKLWSLGQ GIFRGLVNLD RLLLHENQLQ WVHHKAFHDL HRLTTLFLFN HGNRISSVPE RAFRGLHSLD RLLLHQNRVA HVHPHAFRDL GRLMTLYLFA HGNFRGLLD RLLLH-NVHAFL -RLL-LF-
NOGO-R2 NOGO-R3 NOGO-R1 Consensus	NOGO-R2 NOGO-R3 NOGO-R1 Consensus	NOGO-R2 NOGO-R3 NOGO-R1 Consensus	NOGO-R2 NOGO-R3 NOGO-R1 Consensus	NOGO-R2 NOGO-R3 NOGO-R1 Consensus

## FIG. 1B

300 QRARVSSSDV RRFRGSSSAV QKFRGSSSEV R-SSS-V	350  DEEPLGLPKC	400 TLYRDLPA YKKAGKNCTS PRHI.NDSPF	450 QRGEQMCPGA KRKGKCARRT NRTRSHCRLG	I
CRARPLWAWF CRARSLWEWL CRARPLWAWL CRAR-LW-W-	AAPTRPGSRA VSPHQIKSHT YHPIWTGRAT P	AGAPPADPS. HGHPPGSRSG GDSPPGNGSG PPS-	GGED SFDIMPTARP PRRRPGCSRK	49' HL~~~~~~~~ TLR~~~~~~~~
RINANPWACD RINGNAWDCG RINDNPWVCD	FQACP.P FRNCTGP LQGCAVATGP	G.VAE G.ASRDKGHP GNALKGRVPP G	DYWGGY DYAPDYQHKF P.PGFPTSG	LLCLLLLVPH LLAWILGLAV LTCSLTPLGL
LADLPSLEFL LAPLVALEFL LAPLRALQYL LA-LL	RDLRALREAD QDLKLLRVED RDLKRLAAND -DLLD	SSSNH.LY RKEHHPSH VLEPGRPASA	QGGDAPTE.D SSGKELTELQ LTAVRPEGSE	PALSAGLPSP ASSGTALGAP DSEGSGALPS
251 NSLASLPGEA NSLTELQGDC NNLSALPTEA N-LL	301 TCATPPERQG PCATPELRQG PCSLPQRLAG -CPG	351 RGN CQPDAADKAS	401 EDSRGR HRNRNQISKV GTLPGSAEPP	451 ACQAPPDSRG PIRAPSGVQQ QAGSGGGGTG
NOGO-R2 NOGO-R3 NOGO-R1 Consensus	NOGO-R2 NOGO-R3 NOGO-R1 Consensus	NOGO-R2 NOGO-R3 NOGO-R1 Consensus	NOGO-R2 NOGO-R3 NOGO-R1 Consensus	NOGO-R2 NOGO-R3 NOGO-R1 Consensus

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**FIG. 2** 

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	4 FI	6/6 G. 3	
1	PKVTTSCEQQ GLQAVPVGIP AASQRIFLHG NRISHVPAAS FRACRNLTIT PKVTTSCEQQ GLQAVPTGIP ASSQRIFLHG NRISHVPAAS FQSCRNLTIL P.MTVSCQAH NFAAIREGIP EDSERIFLQN NRITFLQQGH FSP. AMVTL ************************************	LER 2  WLHSNVLARI DAAAFTGLAL LEQLDLSDNA QLRSVDPATF HGLGRLHTLH WLHSNALARI DAAAFTGLTL LEQLDLSDNA QLRSVDPATF HGLGRLHTLH WIYSNNITFI APNTFEGFVH LEELDLGDNR QLRTLAPETF QGLVKLHALY WIYSNNITYI HPSTFEGFVH LEELDLGDNR QLRTLAPETF QGLVKLHALY WLFSNNITYI HPSTFEGFVH LEELDLGDNR QLRTLAPETF QGLVKLHALY WLFSNNITYI YPGTFRHLQA LEELDLGDNR HLRSLEPDTF QGLERLQSLH W-SN	LRR 4 200 LDRCGLQELG PGLFRGLAAL QYLYLQDNAL QALPDDTFRD LGNLTHLFLH LDRCGLRELG PGLFRGLAAL QYLYLQDNNL QALPDDTFRD LGNLTHLFLH LYKCGLSALP AGIFGGLHSL QYLYLQDNHI EYLQDDIFVD LVNLSHLFLH LYKCGLSALP AGVFGGLHSL QYLYLQDNHI EYLQDDIFVD LVNLSHLFLH LYKCQLSSLP GNIFRGLYSL QYLYLQENSL LHLQDDLFAD LANLSHLFLH LYCQLSSLP GNIFRGLYSL QYLYLQENSL LHLQDDLFAD LANLSHLFLH LC-LLF-GLL QYLYLQ-NL-DF-D L-NI, HT FT.H
Human NOGO-R1 Murine NOGO-R3 Human NOGO-R3 Human NOGO-R2	Human NOGO-R1 Murine NOGO-R3 Human NOGO-R3 Human NOGO-R2	Human NOGO-R1 Murine NOGO-R1 Murine NOGO-R3 Human NOGO-R3 Human NOGO-R2	Human NOGO-R1 Murine NOGO-R1 Murine NOGO-R3 Human NOGO-R3

LRR 7

LRR 6

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720	RLMTLYLFAN	RLTTEEFNN	RLTTLFLENN	RLTILYLF <b>nn</b>	J RLL-LF- <b>N</b> CT	300	KFRGSSSEVP	KFRGSSSEVP Debassante	RFRGSSSAVE	RARVSSSDVT	R-SSS-V-	350	EEPLGLPKCC	EELLSLPKCC	SDRAARKEHH	TDRAARKEHH	RGNSSSNH	] { } ! ! ! ! ! ! !	400	RHINDSPEGT	RHINDSPEGT	RNRNQISKVS	RNRNQISKAG	EDSRGR	
	VHPHAFRDLG		VHHKAFHDLR	VHR	VHAFL-			RARPEWAMLO			RAR-LW-W		HPIWTGRATD	RPIQTSQLTD	HQIKSHTLTT	HQIKSHTLTT	TRPGSRA	 		DSPPGNGSGP	DTPPGNGSGP	. KAGKNCTSH	. KPGKNCTNP	.DLPA	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
	S LILHQNRVAH			LLLHGN	TTTTI N			L'NGNEWYCDC		LNANPWACDC	LN-N-W-C-C		QGCAVATGPY	EGCAVASGPF			QACP.PAA	- d D		NALKGRVPPG	NALKGRVPPG	SGYK	PGHR	TLYR	           
	AFRGIHSLDR			$\triangleright$	FKGLLUR 8			MFLKSLQYLK APLVALFFTR			LL-		DLKRLAANDL	DLKRLAASDL	DLKLLRVEDF	DLKLLRAEDF	DLRALREADF			LEPGRPASAG	LEPGRPASAG	HPHGHPPGSR	HPHGPR	APPADPS	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
i )	GNRISSVPER	GNKLWSLGQG	GNKLWSLGPG	GNRLRLLTEH	LRR		NESALPTEAL	SLTRIOGDCT.	SLSELQGECL	SLASLPGEAL	TTT	301	CSLPQŘLAGR	CNLPQRLADR	CATPELROGO	CVSPGLRHGQ	$24 ^{\circ}$		351	QPDAADKASV	QPDAADKASV	PSHGASRDKG	SPHGPTRSKG	.LYGVA.EAG	
	Human NOGO-R1 Murine NOGO-R1	Murine NOGO-R3	Human NOGO-R3	Human NOGO-R2			Human NOGO-R1	Murine NOGO-R3	Human NOGO-R3	Human NOGO-R2	Consensus		Human NOGO-R1	Murine NOGO-R1	Murine NOGO-R3	Human NOGO-R3	Human NOGO-KZ	consensus		Human NOGO-R1	Murine NOGO-R1	Murine NOGO-R3	Human NOGO-R3	Human NOGO-R2	Consensus

6,	<b>/</b> 6	
FIG.	3,	cont.

450 RTRSHCRLGQ RTRSHCRLGQ PIRAPSGVQQ PIRAPSGVQQ ACQAPPD	
LPGSAEPPLT AVRPEGSEPP GFPTSGP RRRPGCSRKN RTRSHCRLGQ LPSSAEPPLT ALRPGGSEPP GLPTTGP RRRPGCSRKN RTRSHCRLGQ .SGKELTELQ DYAPDYQHKF SFDIMPTARP KRKGKCARRT PIRAPSGVQQ .AGKQAPELP DYAPDYQHKF SFDIMPTARP KRKGKCARRT PIRAPSGVQQ .QGGDAPTED DYWGGYGGED QRGEQMCPGA ACQAPPD	AGSGGGTGD SEGSGALPSL TCSLTPLGLA LVLWTVLGPC AGSGASGTGD AEGSGALPAL ACSLAPEGLA LVLWTVLGPC
401 LPGSAEPPLT AVR LPSSAEPPLT ALR SGKELTELQ DYA AGKQAPELP DYA QGGDAPTED DYM	451 AGSGGGTGD SEG AGSGASGTGD AEGA
Human NOGO-R1 Murine NOGO-R3 Human NOGO-R3 Human NOGO-R2	Human NOGO-R1 Murine NOGO-R3 Human NOGO-R3 Human NOGO-R2